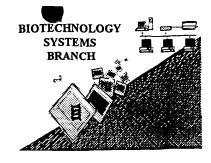
## RAW SEQUENCE LISTING ERROR REPORT



The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 09/873,3/9Source: 0/96Date Processed by STIC: 6/26/2001

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.
PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

- 1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,
- 2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY

FOR CRF SUBMISSION QUESTIONS, PLEASE CONTACT MARK SPENCER, 703-308-4212.

FOR SEQUENCE RULES INTERPRETATION, PLEASE CONTACT ROBERT WAX, 703-308-4216. PATENTIN 2.1 e-mail help: patin21help@uspto.gov or phone 703-306-4119 (R. Wax) PATENTIN 3.0 e-mail help: patin3help@uspto.gov or phone 703-306-4119 (R. Wax)

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE <u>CHECKER</u> <u>VERSION 3.0 PROGRAM</u>, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW:

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The Checker Version 3.0 application is a state-of the-art Windows based software program employing a logical and intuitive user-interface to check whether a sequence listing is in compliance with format and content rules. Checker Version 3.0 works for sequence listings generated for the original version of 37 CFR §§1.821 – 1.825 effective October 1, 1990 (old rules) and the revised version (new rules) effective July 1, 1998 as well as World Intellectual Property Organization (WIPO) Standard ST.25.

Checker Version 3.0 replaces the previous DOS-based version of Checker, and is Y2K-compliant. Checker allows public users to check sequence listings in Computer Readable form (CRF) before submitting them to the United States Patent and Trademark Office (USPTO). Use of Checker prior to filing the sequence listing is expected to result in fewer errored sequence listings, thus saving time and money.

Checker Version 3.0 can be down loaded from the USPTO website at the following address: http://www.uspto.gov/web/offices/pac/checker

OIPE

RAW SEQUENCE LISTING DATE: 06/26/2001 PATENT APPLICATION: US/09/873,319 TIME: 08:49:21

Input Set : N:\jumbos\G15029bk.txt

Output Set: N:\CRF3\06262001\I873319.raw

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3 <110> APPLICANT: Munger, William E.
4 Kulkarni, Prakash
5 Getzenberg, Robert H.
6 Waga, Iwao
7 Yamamoto, Jun
9 <120> TITLE OF INVENTION: Identifying Drugs for and Diagnosis of Benign Prostatic
10 Hyperplasia Using Gene Expression Profiles
12 <130> FILE REFERENCE: 44921-5029-US
14 <140> CURRENT APPLICATION NUMBER: US/09/873,319
15 <141> CURRENT FILING DATE: 2001-06-05
17 <150> PRIOR APPLICATION NUMBER: US 60/223,323
18 <151> PRIOR FILING DATE: 2000-08-07
E--> 20 <160> NUMBER OF SEQ ID NOS: 746
22 <170> SOFTWARE: PatentIn Ver. 2.1
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ERRORED SEQUENCES

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<223> Genbank Accession No. M64594

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I (M)

## Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

VERIFICATION SUMMARY DATE: 06/26/2001 PATENT APPLICATION: US/09/873,319 TIME: 08:49:27

Input Set : N:\jumbos\G15029bk.txt
Output Set: N:\CRF3\06262001\I873319.raw

L:14 M:270 C: Current Application Number differs, Replaced Application Number L:15 M:271 C: Current Filing Date differs, Replaced Current Filing Date L:76 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3 L:92 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4 L:94 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4 L:130 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:6 L:131 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:6 L:151 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7 L:184 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9  $L:185 \ M:341 \ W:$  (46) "n" or "Xaa" used, for SEQ ID#:9 L:188 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9 L:189 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9 L:190 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9 L:228 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:11 L:229 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:11 L:230 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:11 L:264 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:13 L:265 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:13 L:284 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:14 L:301 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:15 L:302 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:15 L:303 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:15 L:304 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:15 L:305 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:15 L:308 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:15 L:329 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:16 L:330 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:16 L:352 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:17 L:353 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:17 L:354 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:17 L:355 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:17 L:393 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:19 L:415 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:20 L:416 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:20 L:417 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:20 L:418 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:20 L:441 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:21 L:492 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:24 L:513 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:25 L:518 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:25 L:519 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:25 L:535 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:26 L:544 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:26 L:567 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:27 L:568 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:27 L:569 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:27 L:585 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:28 L:587 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:28

## VERIFICATION SUMMARY

DATE: 06/26/2001 TIME: 08:49:27 PATENT APPLICATION: US/09/873,319

Input Set : N:\jumbos\G15029bk.txt

Output Set: N:\CRF3\06262001\I873319.raw

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L:20 M:203 E: No. of Seq. differs, <160> Number Of Sequences:Input (746) Counted (748)